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ANALYSIS TOOLS

EBI Generic DB Entry Retrieval

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ID   AE007337    standard; DNA; PRO; 10216 BP.
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DT   31-JUL-2001 (Rel. 68, Created)
DT   02-SEP-2002 (Rel. 72, Last updated, Version 4)
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OC   Bacteria; Firmicutes; Lactobacillales; Streptococcaceae; Streptococcus;
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RX   MEDLINE; 21357209.
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RA   Kolonay J.F., Nelson W.C., Peterson J.D., Umayam L.A., White O.,
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RA   Utterback T.R., Hansen C.L., McDonald L.A., Feldblyum T.V., Angiuoli S.,
RA   Dickinson T., Hickey E.K., Holt I.E., Loftus B.J., Yang F., Smith H.O.,
RA   Venter J.C., Dougherty B.A., Morrison D.A., Hollingshead S.K., Fraser C.M.;
RT   "Complete genome sequence of a virulent isolate of Streptococcus
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RL   Science 293(5529):498-506(2001).
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RA   Venter J.C., Dougherty B.A., Morrison D.A., Hollingshead S.K., Fraser C.M.;
RT   ;
RL   Submitted (29-JUN-2001) to the EMBL/GenBank/DDBJ databases.
RL   The Institute for Genomic Research, 9712 Medical Center Dr, Rockville, MD
RL   20850, USA
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General Description References Links Keywords Sequence

General information

Entry name	**Q97ST2**
Accession number	**Q97ST2**
Created	TrEMBLrel. 18, 1-OCT-2001
Sequence update	TrEMBLrel. 18, 1-OCT-2001
Annotation update	TrEMBLrel. 21, 1-JUN-2002

Description and origin of the Protein

Description	ABC transporter, ATP-binding protein.
Gene name(s)	SP0242.
Organism source	Streptococcus pneumoniae.
Taxonomy	Bacteria; Firmicutes; Bacillus/Clostridium group; Lactobacillales; Streptococcaceae; Streptococcus.
NCBI TaxID	1313

References

[1] Tettelin,H., Nelson,K.E., Paulsen,I.T., Eisen,J.A., Read,T.D., Peterson,S., Heidelberg,J., DeBoy,R.T., Haft,D.H., Dodson,R.J., Durkin,A.S., Gwinn,M., Kolonay,J.F., Nelson,W.C., Peterson,J.D., Umayam,L.A., White,O., Salzberg,S.L., Lewis,M.R., Radune,D., Holtzapple,E., Khouri,H., Wolf,A.M., Utterback,T.R., Hansen,C.L., McDonald,L.A., Feldblyum,T.V., Angiuoli,S., Dickinson,T., Hickey,E.K., Holt,I.E., Loftus,B.J., Yang,F., Smith,H.O., Venter,J.C., Dougherty,B.A., Morrison,D.A., Hollingshead,S.K., Fraser,C.M.,
Complete genome sequence of a virulent isolate of Streptococcus pneumoniae.
(2001) *Science* **293**:498-506

Position	SEQUENCE FROM N.A.
Comments	STRAIN=TIGR4;
Medline	21357209
PubMed	11463916

Database cross-references

EMBL	AE007337; AAK74421.1; -.
TIGR	SP0242; -.
InterPro	IPR003593; AAA_ATPase.
	IPR003439; ABC_transportr.
Pfam	PF00005; ABC_tran; 1.
ProDom	PD000006; ABC_transportr; 1.
SMART	SM00382; AAA; 1.
PROSITE	PS00211; ABC_TRANSPORTER; 1.

Keywords

ATP-binding; Complete proteome;

Sequence information

Length: **363 aa**, molecular weight: **40757 Da**, CRC64 checksum: **3D9F10B1D8DD7731**

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General Description References Links Keywords Sequence